

Serial Number: 08/943,176CRF Processing Date: 1/5/98

Edited by: _____

Verified by: MC (STIC staff)

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☒ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patent bug). Sequences corrected: Seq 2
- ☐ Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/943,776DATE: 01/06/98
TIME: 10:40:55

INPUT SET: S22359.raw

This Raw Listing contains the General
Information Section and up to the first 5 pages.

SEQUENCE LISTING

1
2
3 (1) General Information:
4
5 (i) APPLICANT: Degli-Esposti, Mariapia
6 Goodwin, Raymond
7
8 (ii) TITLE OF INVENTION: Novel Receptor That Causes Cell Death
9
10 (iii) NUMBER OF SEQUENCES: 6
11
12 (iv) CORRESPONDENCE ADDRESS:
13 (A) ADDRESSEE: Immunex
14 (B) STREET: 51 University Street
15 (C) CITY: Seattle
16 (D) STATE: WA
17 (E) COUNTRY: USA
18 (F) ZIP: 98101
19
20 (v) COMPUTER READABLE FORM:
21 (A) MEDIUM TYPE: Floppy disk
22 (B) COMPUTER: Apple Power Macintosh
23 (C) OPERATING SYSTEM: Apple Operating System 7.5.3
24 (D) SOFTWARE: Microsoft Word for Power Macintosh 6.0.1
25
26 (vi) CURRENT APPLICATION DATA:
27 (A) APPLICATION NUMBER:
28 (B) FILING DATE: 03 OCTOBER 1997
29 (C) CLASSIFICATION:
30
31 (vii) PRIOR APPLICATION DATA:
32 (A) APPLICATION NUMBER: USSN 60/044,456
33 (B) FILING DATE: 04 OCTOBER 1996
34 (C) CLASSIFICATION:
35
36 (viii) ATTORNEY/AGENT INFORMATION:
37 (A) NAME: Perkins, Patricia Anne
38 (B) REGISTRATION NUMBER: 34,693
39 (C) REFERENCE/DOCKET NUMBER: 2849-A
40
41 (ix) TELECOMMUNICATION INFORMATION:
42 (A) TELEPHONE: 2065870430
43
44
45 (2) INFORMATION FOR SEQ ID NO:1:
46

RAW SEQUENCE LISTING PATENT APPLICATION US/08/943,776

 DATE: 01/06/98
 TIME: 10:41:01

INPUT SET: S22359.raw

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47      (i) SEQUENCE CHARACTERISTICS:
48          (A) LENGTH: 1847 base pairs
49          (B) TYPE: nucleic acid
50          (C) STRANDEDNESS: single
51          (D) TOPOLOGY: Not Relevant
52
53      (ii) MOLECULE TYPE: cDNA
54
55      (iii) HYPOTHETICAL: NO
56
57      (iv) ANTI-SENSE: NO
58
59      (vii) IMMEDIATE SOURCE:
60          (B) CLONE: AIR
61
62      (ix) FEATURE:
63          (A) NAME/KEY: CDS
64          (B) LOCATION: 236..1489
65
66      (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
67      CTTTTCAGCC ATACCCGGAT GGTTCGTGCC TCGCTGGCCG TGATCACGCC GTCCTCCTTG      60
68
69      GGGATGAGCA GCGCGGCCGT GACGGCGTCC TGGTGCCCCCT CGATCTTGCT CAGCAGCACC      120
70
71      GGGCGGCTGC TCTGCGGCCT GGAGTGGATT TCGGCCGCCA TGTTCGCGCG GCGACTGCTG      180
72
73      CGGCCTCCTC GGCAGGCAGC CCATCAGCTG ACGCCTGGGC GCCCGTCGGA GGGCT ATG      238
74                                     Met
75                                     1
76
77      GAG CAG CGG CCG CGG GGC TGC GCG GCG GTG GCG GCG GCG CTC CTC CTG      286
78      Glu Gln Arg Pro Arg Gly Cys Ala Ala Val Ala Ala Ala Leu Leu Leu
79                                     5                               10                               15
80
81      GTG CTG CTG GGG GCC CGG GCC CAG GGC GGC ACT CGT AGC CCC AGG TGT      334
82      Val Leu Leu Gly Ala Arg Ala Gln Gly Gly Thr Arg Ser Pro Arg Cys
83                                     20                               25                               30
84
85      GAC TGT GCC GGT GAC TTC CAC AAG AAG ATT GGT CTG TTT TGT TGC AGA      382
86      Asp Cys Ala Gly Asp Phe His Lys Lys Ile Gly Leu Phe Cys Cys Arg
87                                     35                               40                               45
88
89      GGC TGC CCA GCG GGG CAC TAC CTG AAG GCC CCT TGC ACG GAG CCC TGC      430
90      Gly Cys Pro Ala Gly His Tyr Leu Lys Ala Pro Cys Thr Glu Pro Cys
91      50                               55                               60                               65
92
93      GGC AAC TCC ACC TGC CTT GTG TGT CCC CAA GAC ACC TTC TTG GCC TGG      478
94      Gly Asn Ser Thr Cys Leu Val Cys Pro Gln Asp Thr Phe Leu Ala Trp
95                                     70                               75                               80
96
97      GAG AAC CAC CAT AAT TCT GAA TGT GCC CGC TGC CAG GCC TGT GAT GAG      526
98      Glu Asn His His Asn Ser Glu Cys Ala Arg Cys Gln Ala Cys Asp Glu
99                                     85                               90                               95
  
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INPUT SET: S22359.raw

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101	CAG	GCC	TCC	CAG	GTG	GCG	CTG	GAG	AAC	TGT	TCA	GCA	GTG	GCC	GAC	ACC		574
102	Gln	Ala	Ser	Gln	Val	Ala	Leu	Glu	Asn	Cys	Ser	Ala	Val	Ala	Asp	Thr		
103			100					105					110					
104																		
105	CGC	TGT	GGC	TGT	AAG	CCA	GGC	TGG	TTT	GTG	GAG	TGC	CAG	GTC	AGC	CAA		622
106	Arg	Cys	Gly	Cys	Lys	Pro	Gly	Trp	Phe	Val	Glu	Cys	Gln	Val	Ser	Gln		
107		115					120					125						
108																		
109	TGT	GTC	AGC	AGT	TCA	CCC	TTC	TAC	TGC	CAA	CCA	TGC	CTA	GAC	TGC	GGG		670
110	Cys	Val	Ser	Ser	Ser	Pro	Phe	Tyr	Cys	Gln	Pro	Cys	Leu	Asp	Cys	Gly		
111	130					135					140					145		
112																		
113	GCC	CTG	CAC	CGC	CAC	ACA	CGG	CTA	CTC	TGT	TCC	CGC	AGA	GAT	ACT	GAC		718
114	Ala	Leu	His	Arg	His	Thr	Arg	Leu	Leu	Cys	Ser	Arg	Arg	Asp	Thr	Asp		
115					150					155					160			
116																		
117	TGT	GGG	ACC	TGC	CTG	CCT	GGC	TTC	TAT	GAA	CAT	GGC	GAT	GGC	TGC	GTG		766
118	Cys	Gly	Thr	Cys	Leu	Pro	Gly	Phe	Tyr	Glu	His	Gly	Asp	Gly	Cys	Val		
119				165					170				175					
120																		
121	TCC	TGC	CCC	ACG	AGC	ACC	CTG	GGG	AGC	TGT	CCA	GAG	CGC	TGT	GCC	GCT		814
122	Ser	Cys	Pro	Thr	Ser	Thr	Leu	Gly	Ser	Cys	Pro	Glu	Arg	Cys	Ala	Ala		
123			180					185					190					
124																		
125	GTC	TGT	GGC	TGG	AGG	CAG	ATG	TTC	TGG	GTC	CAG	GTG	CTC	CTG	GCT	GGC		862
126	Val	Cys	Gly	Trp	Arg	Gln	Met	Phe	Trp	Val	Gln	Val	Leu	Leu	Ala	Gly		
127		195					200					205						
128																		
129	CTT	GTG	GTC	CCC	CTC	CTG	CTT	GGG	GCC	ACC	CTG	ACC	TAC	ACA	TAC	CGC		910
130	Leu	Val	Val	Pro	Leu	Leu	Leu	Gly	Ala	Thr	Leu	Thr	Tyr	Thr	Tyr	Arg		
131	210					215					220					225		
132																		
133	CAC	TGC	TGG	CCT	CAC	AAG	CCC	CTG	GTT	ACT	GCA	GAT	GAA	GCT	GGG	ATG		958
134	His	Cys	Trp	Pro	His	Lys	Pro	Leu	Val	Thr	Ala	Asp	Glu	Ala	Gly	Met		
135					230					235					240			
136																		
137	GAG	GCT	CTG	ACC	CCA	CCA	CCG	GCC	ACC	CAT	CTG	TCA	CCC	TTG	GAC	AGC		1006
138	Glu	Ala	Leu	Thr	Pro	Pro</												

RAW SEQUENCE LISTING PATENT APPLICATION US/08/943,776

DATE: 01/06/98
TIME: 10:41:11

INPUT SET: S22359.raw

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153 AGA GCT CTT GGC CCC GCT GCT GCG CCC ACA CTC TCG CCA GAG TCC CCA 1198
154 Arg Ala Leu Gly Pro Ala Ala Ala Pro Thr Leu Ser Pro Glu Ser Pro
155          310          315          320
156
157 GCC GGC TCG CCA GCC ATG ATG CTG CAG CCG GGC CCG CAG CTC TAC GAC 1246
158 Ala Gly Ser Pro Ala Met Met Leu Gln Pro Gly Pro Gln Leu Tyr Asp
159          325          330          335
160
161 GTG ATG GAC GCG GTC CCA GCG CGG CGC TGG AAG GAG TTC GTG CGC ACG 1294
162 Val Met Asp Ala Val Pro Ala Arg Arg Trp Lys Glu Phe Val Arg Thr
163          340          345          350
164
165 CTG GGG CTG CGC GAG GCA GAG ATC GAA GCC GTG GAG GTG GAG ATC GGC 1342
166 Leu Gly Leu Arg Glu Ala Glu Ile Glu Ala Val Glu Val Glu Ile Gly
167          355          360          365
168
169 CGC TTC CGA GAC CAG CAG TAC GAG ATG CTC AAG CGC TGG CGC CAG CAG 1390
170 Arg Phe Arg Asp Gln Gln Tyr Glu Met Leu Lys Arg Trp Arg Gln Gln
171          370          375          380          385
172
173 CAG CCC GCG GGC CTC GGA GCC GTT TAC GCG GCC CTG GAG CGC ATG GGG 1438
174 Gln Pro Ala Gly Leu Gly Ala Val Tyr Ala Ala Leu Glu Arg Met Gly
175          390          395          400
176
177 CTG GAC GGC TGC GTG GAA GAC TTG CGC AGC CGC CTG CAG CGC GGC CCG 1486
178 Leu Asp Gly Cys Val Glu Asp Leu Arg Ser Arg Leu Gln Arg Gly Pro
179          405          410          415
180
181 TGA CACGGCGCCC ACTTGCCACC TAGGCGCTCT GGTGGCCCTT GCAGAAGCCC 1539
182 *
183
184 TAAGTACGGT TACTTATGCG TGTAGACATT TTATGTCACT TATTAAGCCG CTGGCACGGC 1599
185
186 CCTGCGTAGC AGCACCAGCC GGCCCCACCC CTGCTCGCCC CTATCGCTCC AGCCAAGGCG 1659
187
188 AAGAAGCACG AACGAATGTC GAGAGGGGGT GAAGACATTT CTCAACTTCT CGGCCGGAGT 1719
189
190 TTGGCTGAGA TCGCGGTATT AAATCTGTGA AAGAAAACAA AAAAAAAAAA ACCGGAATTC 1779
191
192 GATATCAAGC TTATCGATAC CGTCGACCTC GAGGGGGGGC CCGGTACCCA ATTCGCCCTA 1839
193
194 TAGTGAGT 1847
195
196
197 (2) INFORMATION FOR SEQ ID NO:2:
198
199 (i) SEQUENCE CHARACTERISTICS:
200 (A) LENGTH: 417 amino acids
201 (B) TYPE: amino acid
202 (D) TOPOLOGY: linear
203
204 (ii) MOLECULE TYPE: protein
205

```

INPUT SET: S22359.raw

	(xi)	SEQUENCE	DESCRIPTION:	SEQ ID NO:2:													
206																	
207																	
208	Met	Glu	Gln	Arg	Pro	Arg	Gly	Cys	Ala	Ala	Val	Ala	Ala	Ala	Leu	Leu	
209	1				5					10					15		
210																	
211	Leu	Val	Leu	Leu	Gly	Ala	Arg	Ala	Gln	Gly	Gly	Thr	Arg	Ser	Pro	Arg	
212				20					25					30			
213																	
214	Cys	Asp	Cys	Ala	Gly	Asp	Phe	His	Lys	Lys	Ile	Gly	Leu	Phe	Cys	Cys	
215			35					40					45				
216																	
217	Arg	Gly	Cys	Pro	Ala	Gly	His	Tyr	Leu	Lys	Ala	Pro	Cys	Thr	Glu	Pro	
218		50					55					60					
219																	
220	Cys	Gly	Asn	Ser	Thr	Cys	Leu	Val	Cys	Pro	Gln	Asp	Thr	Phe	Leu	Ala	
221	65					70					75					80	
222																	
223	Trp	Glu	Asn	His	His	Asn	Ser	Glu	Cys	Ala	Arg	Cys	Gln	Ala	Cys	Asp	
224					85					90					95		
225																	
226	Glu	Gln	Ala	Ser	Gln	Val	Ala	Leu	Glu	Asn	Cys	Ser	Ala	Val	Ala	Asp	
227				100					105					110			
228																	
229	Thr	Arg	Cys	Gly	Cys	Lys	Pro	Gly	Trp	Phe	Val	Glu	Cys	Gln	Val	Ser	
230			115					120					125				
231																	
232	Gln	Cys	Val	Ser	Ser	Ser	Pro	Phe	Tyr	Cys	Gln	Pro	Cys	Leu	Asp	Cys	
233		130					135					140					
234																	
235	Gly	Ala	Leu	His	Arg	His	Thr	Arg	Leu	Leu	Cys	Ser	Arg	Arg	Asp	Thr	
236	145					150					155					160	
237																	
238	Asp	Cys	Gly	Thr	Cys	Leu	Pro	Gly	Phe	Tyr	Glu	His	Gly	Asp	Gly	Cys	
239					165					170					175		
240																	
241	Val	Ser	Cys	Pro	Thr	Ser	Thr	Leu	Gly	Ser	Cys	Pro	Glu	Arg	Cys	Ala	
242				180					185					190			
243																	
244	Ala	Val	Cys	Gly	Trp	Arg	Gln	Met	Phe	Trp	Val	Gln	Val	Leu	Leu	Ala	
245			195					200					205				
246																	
247	Gly	Leu	Val	Val	Pro	Leu											

RAW SEQUENCE LISTING PATENT APPLICATION US/08/943,776

DATE: 01/06/98
TIME: 10:41:22

INPUT SET: S22359.raw

***** PREVIOUSLY ERRORED SEQUENCES - EDITED *****

```

197 (2) INFORMATION FOR SEQ ID NO:2:
198
199 (i) SEQUENCE CHARACTERISTICS:
200 (A) LENGTH: 417 amino acids
201 (B) TYPE: amino acid
202 (D) TOPOLOGY: linear
203
204 (ii) MOLECULE TYPE: protein
205
206 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
207
208 Met Glu Gln Arg Pro Arg Gly Cys Ala Ala Val Ala Ala Ala Leu Leu
209 1 5 10 15
210
211 Leu Val Leu Leu Gly Ala Arg Ala Gln Gly Gly Thr Arg Ser Pro Arg
212 20 25 30
213
214 Cys Asp Cys Ala Gly Asp Phe His Lys Lys Ile Gly Leu Phe Cys Cys
215 35 40 45
216
217 Arg Gly Cys Pro Ala Gly His Tyr Leu Lys Ala Pro Cys Thr Glu Pro
218 50 55 60
219
220 Cys Gly Asn Ser Thr Cys Leu Val Cys Pro Gln Asp Thr Phe Leu Ala
221 65 70 75 80
222
223 Trp Glu Asn His His Asn Ser Glu Cys Ala Arg Cys Gln Ala Cys Asp
224 85 90 95
225
226 Glu Gln Ala Ser Gln Val Ala Leu Glu Asn Cys Ser Ala Val Ala Asp
227 100 105 110
228
229 Thr Arg Cys Gly Cys Lys Pro Gly Trp Phe Val Glu Cys Gln Val Ser
230 115 120 125
231
232 Gln Cys Val Ser Ser Ser Pro Phe Tyr Cys Gln Pro Cys Leu Asp Cys
233 130 135 140
234
235 Gly Ala Leu His Arg His Thr Arg Leu Leu Cys Ser Arg Arg Asp Thr
236 145 150 155 160
237
238 Asp Cys Gly Thr Cys Leu Pro Gly Phe Tyr Glu His Gly Asp Gly Cys
239 165 170 175
240
241 Val Ser Cys Pro Thr Ser Thr Leu Gly Ser Cys Pro Glu Arg Cys Ala
242 180 185 190
243

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DATE: 01/06/98
TIME: 10:41:27

INPUT SET: S22359.raw[illegible]

PAGE: 1

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/08/943,776

DATE: 01/06/98
TIME: 10:41:31

INPUT SET: S22359.raw

Line

Error

Original Text

W. Lazar

1812

PAGE: 1

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/943,776

DATE: 01/05/98
TIME: 14:55:49

INPUT SET: S22359.raw

This Raw Listing contains
Information Section and
containing ERRORS.

Does Not Comply
Corrected Diskette Needed

SEQUENCE LISTING

1
2
3 (1) General Information:
4
5 (i) APPLICANT: Degli-Esposti, Mariapia
6 Goodwin, Raymond
7
8 (ii) TITLE OF INVENTION: Novel Receptor That Causes Cell Death
9
10 (iii) NUMBER OF SEQUENCES: 6
11
12 (iv) CORRESPONDENCE ADDRESS:
13 (A) ADDRESSEE: Immunex
14 (B) STREET: 51 University Street
15 (C) CITY: Seattle
16 (D) STATE: WA
17 (E) COUNTRY: USA
18 (F) ZIP: 98101
19
20 (v) COMPUTER READABLE FORM:
21 (A) MEDIUM TYPE: Floppy disk
22 (B) COMPUTER: Apple Power Macintosh
23 (C) OPERATING SYSTEM: Apple Operating System 7.5.3
24 (D) SOFTWARE: Microsoft Word for Power Macintosh 6.0.1
25
26 (vi) CURRENT APPLICATION DATA:
27 (A) APPLICATION NUMBER:
28 (B) FILING DATE: 03 OCTOBER 1997
29 (C) CLASSIFICATION:
30
31 (vii) PRIOR APPLICATION DATA:
32 (A) APPLICATION NUMBER: USSN 60/044,456
33 (B) FILING DATE: 04 OCTOBER 1996
34 (C) CLASSIFICATION:
35
36 (viii) ATTORNEY/AGENT INFORMATION:
37 (A) NAME: Perkins, Patricia Anne
38 (B) REGISTRATION NUMBER: 34,693
39 (C) REFERENCE/DOCKET NUMBER: 2849-A
40
41 (ix) TELECOMMUNICATION INFORMATION:
42 (A) TELEPHONE: 2065870430
43
44

ERRORED SEQUENCES FOLLOW:

(2) INFORMATION FOR SEQ ID NO:2:

08/943,776

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 417 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

next page

(ii) MOLECULE TYPE: protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Met	Glu	Gln	Arg	Pro	Arg	Gly	Cys	Ala	Ala	Val	Ala	Ala	Ala	Leu	Leu
1				5					10					15	
Leu	Val	Leu	Leu	Gly	Ala	Arg	Ala	Gln	Gly	Gly	Thr	Arg	Ser	Pro	Arg
			20					25					30		
Cys	Asp	Cys	Ala	Gly	Asp	Phe	His	Lys	Lys	Ile	Gly	Leu	Phe	Cys	Cys
		35					40					45			
Arg	Gly	Cys	Pro	Ala	Gly	His	Tyr	Leu	Lys	Ala	Pro	Cys	Thr	Glu	Pro
	50					55					60				
Cys	Gly	Asn	Ser	Thr	Cys	Leu	Val	Cys	Pro	Gln	Asp	Thr	Phe	Leu	Ala
65					70					75					80
Trp	Glu	Asn	His	His	Asn	Ser	Glu	Cys	Ala	Arg	Cys	Gln	Ala	Cys	Asp
				85					90					95	
Glu	Gln	Ala	Ser	Gln	Val	Ala	Leu	Glu	Asn	Cys	Ser	Ala	Val	Ala	Asp
			100					105					110		
Thr	Arg	Cys	Gly	Cys	Lys	Pro	Gly	Trp	Phe	Val	Glu	Cys	Gln	Val	Ser
		115					120					125			
Gln	Cys	Val	Ser	Ser	Ser	Pro	Phe	Tyr	Cys	Gln	Pro	Cys	Leu	Asp	Cys
		130				135					140				
Gly	Ala	Leu	His	Arg	His	Thr	Arg	Leu	Leu	Cys	Ser	Arg	Arg	Asp	Thr
145					150					155					160
Asp	Cys	Gly	Thr	Cys	Leu	Pro	Gly	Phe	Tyr	Glu	His	Gly	Asp	Gly	Cys
				165					170					175	
Val	Ser	Cys	Pro	Thr	Ser	Thr	Leu	Gly	Ser	Cys	Pro	Glu	Arg	Cys	Ala
			180					185					190		
Ala	Val	Cys	Gly	Trp	Arg	Gln	Met	Phe	Trp	Val	Gln	Val	Leu	Leu	Ala
		195					200					205			
Gly	Leu	Val	Val	Pro	Leu	Leu	Leu	Gly	Ala	Thr	Leu	Thr	Tyr	Thr	Tyr
	210					215					220				
Arg	His	Cys	Trp	Pro	His	Lys	Pro	Leu	Val	Thr	Ala	Asp	Glu	Ala	Gly
225					230					235					240
Met	Glu	Ala	Leu	Thr	Pro	Pro	Pro	Ala	Thr	His	Leu	Ser	Pro	Leu	Asp
				245					250					255	
Ser	Ala	His	Thr	Leu	Leu	Ala	Pro	Pro	Asp	Ser	Ser	Glu	Lys	Ile	Cys

02/943,776

260

265

270

Thr Val Gln Leu Val Gly Asn Ser Trp Thr Pro Gly Tyr Pro Glu Thr
 275 280 285

Gln Glu Ala Leu Cys Pro Gln Val Thr Trp Ser Trp Asp Gln Leu Pro
 290 295 300

Ser Arg Ala Leu Gly Pro Ala Ala Ala Pro Thr Leu Ser Pro Glu Ser
 305 310 315 320

Pro Ala Gly Ser Pro Ala Met Met Leu Gln Pro Gly Pro Gln Leu Tyr
 325 330 335

Asp Val Met Asp Ala Val Pro Ala Arg Arg Trp Lys Glu Phe Val Arg
 340 345 350

Thr Leu Gly Leu Arg Glu Ala Glu Ile Glu Ala Val Glu Val Glu Ile
 355 360 365

Gly Arg Phe Arg Asp Gln Gln Tyr Glu Met Leu Lys Arg Trp Arg Gln
 370 375 380

Gln Gln Pro Ala Gly Leu Gly Ala Val Tyr Ala Ala Leu Glu Arg Met
 385 390 395 400

Gly Leu Asp Gly Cys Val Glu Asp Leu Arg Ser Arg Leu Gln Arg Gly
 405 410 415

Pro

*Delete